

Amendments to the Claims:

This listing of claims will replace all prior versions, and listings, of claims in the application:

1.-102. (Canceled).

103. (Currently Amended) A method of inhibiting programmed cell death in a plant or yeast eukaryote, said method comprising:

administering to the plant or yeast eukaryote a bacterial effector protein which inhibits programmed cell death, wherein the protein comprises either: (1) the amino acid sequence of SEQ ID NO: 2; (2) ~~an the amino acid motif sequence spanning a C terminus of SEQ[[.]] ID[[.]] NO[[.]]: 2 selected from the group consisting of the motif of SEQ ID NO: 9, the motif of SEQ ID NO: 10, the motif of SEQ ID NO: 11, the motif of SEQ ID NO: 12, the motif of SEQ ID NO: 13, the motif of SEQ ID NO: 14, the motif of SEQ ID NO: 15, the motif of SEQ ID NO:16, the motif of SEQ ID NO: 17, the motif of SEQ ID NO: 18, the motif of SEQ ID NO: 19, the motif of SEQ ID NO:20, the motif of SEQ ID NO: 21, the motif of SEQ ID NO: 22, the motif of SEQ ID NO: 23, and combinations thereof; or (3) the amino acid sequence spanning amino acids 308 and 553 of SEQ[[.]] ID[[.]] NO[[.]]: 2.~~

104. (Currently Amended) The method according to claim 103, wherein the protein comprises the amino acid sequence of SEQ[[.]] ID[[.]] NO[[.]]: 2.

105.-106. (Canceled).

107. (Currently Amended) The method according to claim 103, wherein the protein comprises the amino acid ~~motif sequence spanning a C terminus of SEQ[[.]] ID[[.]] NO[[.]]: 2 selected from the group consisting of the motif of SEQ ID NO: 9, the motif of SEQ ID NO: 10, the motif of SEQ ID NO: 11, the motif of SEQ ID NO: 12, the motif of SEQ ID NO: 13, the motif of SEQ ID NO: 14, the motif of SEQ ID NO: 15, the motif of SEQ ID NO:16, the motif of SEQ ID NO: 17, the motif of SEQ ID NO: 18, the motif of SEQ ID NO: 19, the motif of SEQ ID NO:20, the motif of SEQ ID NO: 21, the motif of SEQ ID NO: 22, the motif of SEQ ID NO: 23, and combinations thereof.~~

108. (Currently Amended) The method according to claim 103, wherein the protein comprises the amino acid sequence spanning amino acids 308 and 553 of SEQ[[.]] ID[[.]] NO[[.]]: 2.

109. (Canceled).

110. (Previously Presented) The method according to claim 103, wherein the eukaryote is a plant.

111. (Currently Amended) The method according to claim 103, wherein the eukaryote is a plant yeast.